

Mark N Puttick

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

2,741
citations

331259

21
h-index

377514

34
g-index

36
all docs

36
docs citations

36
times ranked

3707
citing authors

#	ARTICLE	IF	CITATIONS
1	Coevolution of enamel, ganoin, enameloid, and their matrix SPCP genes in osteichthyans. <i>IScience</i> , 2021, 24, 102023.	1.9	27
2	Empirical distributions of homoplasy in morphological data. <i>Palaeontology</i> , 2021, 64, 505-518.	1.0	9
3	Phylogenetic sampling affects evolutionary patterns of morphological disparity. <i>Palaeontology</i> , 2021, 64, 765-787.	1.0	6
4	MOTMOT: Models of trait macroevolution on trees (an update). <i>Methods in Ecology and Evolution</i> , 2020, 11, 464-471.	2.2	14
5	Arachnid monophyly: Morphological, palaeontological and molecular support for a single terrestrialization within Chelicerata. <i>Arthropod Structure and Development</i> , 2020, 59, 100997.	0.8	35
6	The complex effects of mass extinctions on morphological disparity. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 2207-2220.	1.1	19
7	Shifting spaces: Which disparity or dissimilarity measurement best summarize occupancy in multidimensional spaces?. <i>Ecology and Evolution</i> , 2020, 10, 7261-7275.	0.8	54
8	A Cambrian–Ordovician Terrestrialization of Arachnids. <i>Frontiers in Genetics</i> , 2020, 11, 182.	1.1	43
9	Disparities in the analysis of morphological disparity. <i>Biology Letters</i> , 2020, 16, 20200199.	1.0	60
10	Probabilistic methods outperform parsimony in the phylogenetic analysis of data simulated without a probabilistic model. <i>Palaeontology</i> , 2019, 62, 1-17.	1.0	44
11	MCMCtreeR: functions to prepare MCMCtree analyses and visualize posterior ages on trees. <i>Bioinformatics</i> , 2019, 35, 5321-5322.	1.8	128
12	Characterization of melanosomes involved in the production of non-iridescent structural feather colours and their detection in the fossil record. <i>Journal of the Royal Society Interface</i> , 2019, 16, 20180921.	1.5	17
13	Origin of horsetails and the role of whole-genome duplication in plant macroevolution. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191662.	1.2	17
14	Archosauromorph extinction selectivity during the Triassic–Jurassic mass extinction. <i>Palaeontology</i> , 2019, 62, 211-224.	1.0	20
15	The timescale of early land plant evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E2274-E2283.	3.3	654
16	Empirical realism of simulated data is more important than the model used to generate it: a reply to Goloboff <i>et al.</i> . <i>Palaeontology</i> , 2018, 61, 631-635.	1.0	29
17	The Interrelationships of Land Plants and the Nature of the Ancestral Embryophyte. <i>Current Biology</i> , 2018, 28, 733-745.e2.	1.8	398
18	Mixed evidence for early bursts of morphological evolution in extant clades. <i>Journal of Evolutionary Biology</i> , 2018, 31, 502-515.	0.8	28

#	ARTICLE	IF	CITATIONS
19	Probabilistic methods surpass parsimony when assessing clade support in phylogenetic analyses of discrete morphological data. <i>Palaeontology</i> , 2018, 61, 105-118.	1.0	61
20	Evolution of jaw disparity in fishes. <i>Palaeontology</i> , 2018, 61, 847-854.	1.0	21
21	Reply to Hedges et al.: Accurate timetrees do indeed require accurate calibrations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E9512-E9513.	3.3	15
22	Evolution of metazoan morphological disparity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E8909-E8918.	3.3	78
23	Well-Annotated microRNAomes Do Not Evidence Pervasive miRNA Loss. <i>Genome Biology and Evolution</i> , 2018, 10, 1457-1470.	1.1	41
24	Integrated genomic and fossil evidence illuminates life's early evolution and eukaryote origin. <i>Nature Ecology and Evolution</i> , 2018, 2, 1556-1562.	3.4	274
25	Uncertain-tree: discriminating among competing approaches to the phylogenetic analysis of phenotype data. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20162290.	1.2	114
26	Parsimony and maximum-likelihood phylogenetic analyses of morphology do not generally integrate uncertainty in inferring evolutionary history: a response to Brown et al. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20171636.	1.2	19
27	Body length of bony fishes was not a selective factor during the biggest mass extinction of all time. <i>Palaeontology</i> , 2017, 60, 727-741.	1.0	13
28	Bayesian methods outperform parsimony but at the expense of precision in the estimation of phylogeny from discrete morphological data. <i>Biology Letters</i> , 2016, 12, 20160081.	1.0	160
29	Dating placentalia: Morphological clocks fail to close the molecular fossil gap. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 873-886.	1.1	26
30	Partially incorrect fossil data augment analyses of discrete trait evolution in living species. <i>Biology Letters</i> , 2016, 12, 20160392.	1.0	30
31	A molecular palaeobiological exploration of arthropod terrestrialization. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150133.	1.8	131
32	Fossils and living taxa agree on patterns of body mass evolution: a case study with Afrotheria. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20152023.	1.2	27
33	Size is not everything: rates of genome size evolution, not C -value, correlate with speciation in angiosperms. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20152289.	1.2	65
34	HIGH RATES OF EVOLUTION PRECEDED THE ORIGIN OF BIRDS. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 1497-1510.	1.1	63